Resource Summary Report

Generated by PRECISE-TBI on Jul 31, 2026

Hypothes.is

RRID:SCR_000430
Type: Tool

Proper Citation

Hypothes.is (RRID:SCR_000430)

Resource Information

URL: http://hypothes.is/

Proper Citation: Hypothes.is (RRID:SCR_000430)

Description: Non-profit organization working towards developing an open platform for the collaborative evaluation of knowledge. It will combine sentence-level critique with community peer-review to provide commentary, references, and insight on top of news, blogs, scientific articles, books, terms of service, ballot initiatives, legislation and regulations, software code and more.

Abbreviations: Hypothes.is

Resource Type: software resource

Keywords: annotation, evaluation, annotation standard, standard specification, peer review, ASWG

Funding: Alfred P. Sloan Foundation ;
Shuttleworth Foundation ;
Andrew W. Mellon Foundation

Resource Name: Hypothes.is

Resource ID: SCR_000430

Alternate IDs: nlx_156054

Record Creation Time: 20220129T080201+0000

Record Last Update: 20260725T190444+0000

Ratings and Alerts

No rating or validation information has been found for Hypothes.is.

No alerts have been found for Hypothes.is.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 11 mentions in open access literature.

Listed below are recent publications. The full list is available at PRECISE-TBI .

Piekniewska A, et al. (2025) Do organisms need an impact factor? Citations of key biological resources including model organisms reveal usage patterns and impact. PloS one, 20(8), e0327344.

Bandrowski A, et al. (2023) The Antibody Registry: ten years of registering antibodies. Nucleic acids research, 51(D1), D358.

Riedel N, et al. (2022) Replacing bar graphs of continuous data with more informative graphics: are we making progress? Clinical science (London, England : 1979), 136(15), 1139.

Weissgerber T, et al. (2021) Automated screening of COVID-19 preprints: can we help authors to improve transparency and reproducibility? Nature medicine, 27(1), 6.

Hsu CN, et al. (2020) Comparing the Use of Research Resource Identifiers and Natural Language Processing for Citation of Databases, Software, and Other Digital Artifacts. Computing in science & engineering, 22(2), 22.

Brito JJ, et al. (2020) Recommendations to enhance rigor and reproducibility in biomedical research. GigaScience, 9(6).

Babic Z, et al. (2019) Incidences of problematic cell lines are lower in papers that use RRIDs to identify cell lines. eLife, 8.

Erwin T, et al. (2018) ZooKeys anniversary: 10 years of leadership toward open-access publishing of zoological data and establishment at Pensoft of like-minded sister journals across the biodiversity spectrum. ZooKeys(770), 1.

Kress WJ, et al. (2018) PhytoKeys at 100: progress in sustainability, innovation, and speed to enhance publication in plant systematics. PhytoKeys(100), 1.

Garcia A, et al. (2018) Biotea: semantics for Pubmed Central. PeerJ, 6, e4201.

Wilson G, et al. (2014) Software Carpentry: lessons learned. F1000Research, 3, 62.